

**AMENDMENTS TO THE CLAIMS**

This listing of claims will replace all prior versions, and listings, of claims in the application:

**Listing of Claims:**

1. (Original) A method of multiple microorganism detection which is a method for detecting two or more microorganisms having different properties in foods, with high sensitivity comparable or even superior to official methods, by amplifying a plurality of target genes with a single PCR reaction tube and analyzing the same, comprising the following steps:
  - (A) a step for extracting DNA of the target microorganisms to be detected, by treating at least with a lytic enzyme and/or bacteriocin having lytic activity, a surfactant and a protein denaturant; and
  - (B) a step for performing Multiplex PCR by mixing a primer specific to the target microorganisms to be detected.
2. (Original) The method of multiple microorganism detection according to claim 1, wherein a step to culture microorganisms under a culture condition where 1 CFU/100 g microorganisms become  $10^3$  CFU/ml or more after 24 h of culture, is included prior to the step of extracting DNA of the target microorganisms to be detected.
3. (Original) The method of multiple microorganism detection according to claim 1 or 2, wherein the two or more microorganisms with different properties comprise *Listeria monocytogenes*.
4. (Original) The method of multiple microorganism detection according to claim 3, wherein the specific primer is a primer consisting of base sequences shown by SEQ ID Nos: 5 and 6.

5. (Withdrawn) The method of multiple microorganism detection according to claim 1 or 2, wherein the two or more microorganisms with different properties comprise pathogenic *Escherichia coli* O157.
6. (Withdrawn) The method of multiple microorganism detection according to claim 5, wherein the specific primer is a primer consisting of base sequences shown by SEQ ID Nos: 1 and 2, or SEQ ID Nos: 7 and 8.
7. (Withdrawn) The method of multiple microorganism detection according to claim 1 or 2, wherein the two or more microorganisms with different properties comprise *Salmonella* spp.
8. (Withdrawn) The method of multiple microorganism detection according to claim 7, wherein the specific primer is a primer consisting of base sequences shown by SEQ ID Nos: 3 and 4, or SEQ ID Nos: 9 and 10.
9. (Currently amended) The method of multiple microorganism detection according to ~~any one of claims 1 to 8~~ claim 1 or 2, wherein the microorganisms are cultured in a culture condition where the pH after culture becomes 5.1 or more.
10. (Currently amended) The method of multiple microorganism detection according to ~~any one of claims 1 to 9~~ claim 1 or 2, wherein the microorganisms are cultured in a medium with glucose concentration of 0.15% or less, and/or in a medium with concentration of phosphate-buffer solution of 50 mM or more or in a medium with a buffer ability similar as that with concentration of phosphate-buffer solution of 50 mM or more.
11. (Currently amended) The method of multiple microorganism detection according to ~~any one of claims 1 to 10~~ claim 1 or 2, wherein the extraction is performed after treating with a lytic enzyme and/or bacteriosin having a lytic activity, further treating with a surfactant and a protein denaturant, removing insoluble fractions by centrifugation, and by depositing DNA by alcohol precipitation.

12. (Currently amended) The method of multiple microorganism detection according to ~~any one of claims 1 to 14~~ claim 1 or 2, wherein the lytic enzyme is Achromopeptidase and/or lysozyme.
13. (Currently amended) The method of multiple microorganism detection according to ~~any one of claims 1 to 12~~ claim 1 or 2, wherein bacteriosin having lytic activity is Enterolysine.
14. (Currently amended) The method of multiple microorganism detection according to ~~any one of claims 1 to 13~~ claim 1 or 2, wherein the surfactant is ethyleneoxide condensate of sorbitan monolaurate.
15. (Currently amended) The method of multiple microorganism detection according to ~~any one of claims 1 to 14~~ claim 1 or 2, wherein the protein denaturant is Guanidine isothiocyanate.
16. (Currently amended) The method of multiple microorganism detection according to ~~any one of claims 1 to 15~~ claim 1 or 2, wherein Multiplex PCR is performed by combining DNA consisting of base sequences shown by SEQ ID NOs: 1 to 6 at a total concentration of 750 nM or less as a primer.
17. (Withdrawn- currently amended) The method of multiple microorganism detection according to ~~any one of claims 1 to 15~~ claim 1 or 2, wherein Multiplex PCR is performed by combining DNA consisting of base sequences shown by SEQ ID NOs: 5 to 10 at a total concentration of 750 nM or less as a primer.
18. (Currently amended) The method of multiple microorganism detection according to ~~any one of claims 1 to 17~~ claim 1 or 2, wherein the food is edible meat or processed meat product.